## Claims

- 1. Method for detection of endotoxin, comprising the steps:
  - a) incubation of a sample with a bacteriophage tail protein, and subsequently
  - b) detection of endotoxin bound to bacteriophage tail proteins by means of spectroscopic methods, ELISA, chemical or enzymatic detection reaction of endotoxins or cleaved-off endotoxin components, or by means of capacitance measurements..
- 2. Method according to claim 1, further comprising after step a) and prior to step b) the additional step
  - a') separation of the bacteriophage tail protein-endotoxin complexes from the sample.
- 3. Method for detection of endotoxin, the method comprising the steps of:
  - a) contacting a sample containing endotoxins with a surface, subsequently
  - b) incubating of bacteriophage tail proteins with the endotoxin immobilised on the surface, and
  - c) detection of bacteriophage tail proteins by means of spectroscopic methods, ELISA, chemical or enzymatic detection reaction of endotoxins or cleaved-off endotoxin components, or by means of capacitance measurements.
- 4. Method according to claim 3 further comprising after step b) and before step c) an additional step
  - b') separation of said bound bacteriophage tail proteins from endotoxin.
- 5. Method according to any one of the preceding claims, wherein the bacteriophage tail protein is a protein of the short bacteriophage tail fiber or a coat protein of bacteriophages without tail.

- 6. Method according to claim 5, wherein the protein of the short bacteriophage tail fiber is selected from K3, T2, T4, Ox2, RB32-33, AR1, PP01 and RB69.
- 7. Method according to claims 5 or 6, wherein the bacteriophage tail protein has a homology of at least 60 % to T4p12 protein on the amino acid level.
- 8. Method according to any one of the preceding claims, wherein the bacteriophage tail proteins are modified.
- 9. Method according to any one of the preceding claims, wherein the bacteriophage tail proteins are covalently linked to enzymatically active proteins.
- 10. Method according to any one of the preceding claims, wherein the bacteriophage tail protein comprises a strep-tag or a his-tag.
- 11. Method according to claim 10, wherein the tag comprises an amino acid sequence according to SEQ ID NOS 5, 6 or 7.
- 12. Method according to claim 10 or 11, wherein the p12-protein of phage T4, K3, T2, Ox2, RB32-33, AR1, PP01 or RB69 is used as bacteriophage tail protein.
- 13. Method according to any one of the preceding claims, wherein the  $Ca^{2+}$  concentration is in the incubation 0.1  $\mu$ M to 10 mM and/or the  $Mg^{2+}$  concentration is 0.1  $\mu$ M to 10 mM.
- 14. Method according to any one of the preceding claims, wherein marked endotoxin is displaced from the binding with a bacteriophage tail protein and wherein the marked endotoxin is detected subsequently.
- 15. An endotoxin detection kit comprising a carrier coated with bacteriophage tail proteins, a container containing a reference endotoxin for measurement of a standard curve, a container with at least one further bacteriophage tail protein or an anti lipid A antibody.